

SEQUENCE LISTING

Keller, Richard Werner, James Goodwin, Peter <120> RAPID HAPLOTYPING BY SINGLE MOLECULE DETECTION <130> S-94,652 <140> US 09/862,855 <141> 2003-04-11 <160> 21 <170> PatentIn version 3.1 <210> 1 <211> 20 <212> DNA <213> Unknown <220> Feature: <223> 20-mer probe complementary to 20 base pair region of M13mp18 target containing EcoR I restriction site <400> 1 20 gctcgaattc gtaatcatcg <210> 2 <211> 18 <212> DNA <213> Unknown <220> Feature: <223> 18-mer probe complementary to 18 base pair region of M13mp18 target containing Hind III restriction site <400> 2 18 cagtgccaag cttcgatg <210> 3 <211> 97 <212> DNA <213> Unknown <220> Feature: <223> Synthetic chimera template derived from genes MLL(HRX, Htrx) and AF4(FEL) gaagttccca aaaccactcc tagtgagccc aagaaaaagc agcctccacc accaaaacaa 60 97 tatgatacat cttcaaaaac tcactcaaat tctcagc

<210> 4 <211> 27

```
<212> DNA
<213> Unknown
<220> Feature:
<223> DNA probe sequence complementary to MLL portion of SEQ ID NO:3
<400> 4
aaaaatttct tgggcttcac tagggag
                                                                     27
<210> 5
<211> 29
<212> DNA
<213> Unknown
<220> Feature:
<223> DNA probe sequence complementary to AF4 portion of SEQ ID NO:3
<400> 5
aaaaaaattt gagtgagttt ttgaagatg
                                                                     29
<210> 6
<211> 12
<212> DNA
<213> Unknown
<220> Feature:
<223> PNA probe sequence for MLL portion of SEQ ID NO:3
<400> 6
tttcttgggc tc
                                                                     12
<210> 7
<211> 12
<212> DNA
<213> Unknown
<220> Feature:
<223> PNA probe sequence for AF4 portion of SEQ ID NO:3
<400> 7
tttgagtgag tt
                                                                     12
<210> 8
<211> 12
<212> DNA
<213> Unknown
<220> Feature:
<223> LNA probe sequence for MLL portion of SEQ ID NO:3
<400> 8
tttcttgggc tc
                                                                     12
<210> 9
<211> 12
<212> DNA
```

<213> Unknown

```
<220> Feature:
<223> LNA probe sequence for AF4 portion of SEQ ID NO:3
<400> 9
tttgagtgag tt
                                                                      12
<210> 10
<211> 32
<212> DNA
<213> Unknown
<220> Feature:
<223> Synthetic oligonucleotide template containing a subset of a sequence
variant for the HLA gene
<400> 10
tggcagctca gaccaccaag cacaagtggg ag
                                                                      32
<210> 11
<211> 76
<212> DNA
<213> Unknown
<220> Feature:
<223> Synthetic oligonucleotide template containing a subset of a sequence
variant for the HLA gene
<400> 11
geggeeeatg tggeggagea gttgagagee tacetggagg geaegtgegt ggagtggete
                                                                      60
cgcagatacc tggaga
                                                                      76
<210> 12
<211> 32
<212> DNA
<213> Unknown
<220> Feature:
<223> Synthetic oligonucleotide template containing a subset of a sequence
variant for the HLA gene
<400> 12
tggcagctca gaccaccaag cacaagtggg ag
                                                                      32
<210> 13
<211> 76
<212> DNA
<213> Unknown
<220> Feature:
<223> Synthetic oligonucleotide template containing a subset of a sequence
variant for the HLA gene
gcggcccatg tggcggagca gcagagagcc tacctggagg gcacgtgcgt ggagtggctc
                                                                     60
cgcagatacc tggaga
                                                                     76
```

| <210> 14 | |
|--|---------|
| <211> 32 | |
| <212> DNA | |
| <213> Unknown | |
| <pre><220> Feature: </pre> | uence |
| <223> Synthetic oligonucleotide template containing a base. | |
| variant for the HLA gene | |
| | |
| <400> 14 | 32 |
| tggcagctca gaccacccaa gacaagtggg ag | |
| | |
| | |
| <210> 15 | |
| <211> 76 | |
| <212> DNA | |
| <213> Unknown | |
| <pre><220> Feature: <223> Synthetic oligonucleotide template containing a subset of a sec</pre> | quence |
| <223> Synthetic oligonucleotide template containing a summer | |
| variant for the HLA gene | |
| | |
| <400> 15 Language gracet grace | 60 |
| <400> 15 gcggcccatg tggcggagca gttgagagcc tacctggagg gcacgtgcgt ggacgggctc | |
| | 76 |
| cgcagatacc tggaga | |
| | |
| | |
| <210> 16 | |
| <211> 32 | |
| <212> DNA | |
| <213> Unknown | |
| <pre><220> Feature: <223> Synthetic oligonucleotide template containing a subset of a se</pre> | equence |
| <223> Synthetic oligonacieociae compens | |
| variant for the HLA gene | |
| | |
| <400> 16 | 32 |
| tggcagctca gaccacccaa ggcaagtggg ag | |
| | |
| 010. 17 | |
| <210> 17 <211> 76 | |
| 1= | |
| <212> DNA <213> Unknown | |
| | canence |
| <220> Feature: <223> Synthetic oligonucleotide template containing a subset of a s | equence |
| variant for the HLA gene | |
| Variance for the second | |
| <400> 17 | 60 |
| <400> 17 gcggcccatg tggcggagca gcagagagcc tacctggagg gcacgtgcgt ggacgggctc | |
| 3-33 | 76 |
| cgcagatacc tggaga | · - |
| | |
| | |
| <210> 18 | |
| <211> 32 | |
| <212> DNA | |

| <220> <223> | - | template containing a | subset of a | sequence |
|-------------------------------------|---|-----------------------|-------------|----------|
| <400> tggcago | 18 etca gaccacccaa ggcaagtggg | ag | | 32 |
| | | | | |
| | Synthetic oligonucleotide for the HLA gene | template containing a | subset of a | sequence |
| <400> gcggccc | 19 atg tggcggagca gcagagagcc | tacctggagg gcacgtgcgt | ggagtggctc | 60 |
| cgcagat | acc tggaga | | | 76 |
| <212><213><220><223> | 20 32 DNA Unknown Feature: Synthetic oligonucleotide for the HLA gene | template containing a | subset of a | sequence |
| | 20 tca gaccacccaa ggcaagtggg | ag | | 32 |
| <210><211><212><212><213><220><223> | 21 76 | | subset of a | sequence |
| | 21 atg tggcggagca gttgagagcc | tacctggagg gcacgtgcgt | ggacgggctc | 60 |

cgcagatacc tggaga

76